Amendment to the claims

This listing of claims replaces all prior versions of claims in the application:

Listing of Claims

- 1-32 (cancelled)
- 33. (currently amended) An isolated polypeptide having at least 80% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[)]; or
- b) the amino acid sequence of the polypeptide shown in Figure 2 (SEQ ID NO:2), lacking its associated signal peptide; or
- [(c)](b) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
- 34. (currently amended) The isolated polypeptide of Claim 33 having at least 85% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[[)]]; or
- (b) the amino acid sequence of the polypeptide shown in Figure 2 (SEQ ID NO:2), lacking its associated signal peptide; or
- [(c)](b) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
- 35. (currently amended) The isolated polypeptide of Claim 33 having at least 90% amino acid sequence identity to:

- (a) the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[[)]; or
- (b) the amino acid sequence of the polypeptide shown in Figure 2 (SEQ ID NO:2), lacking its associated signal peptide; or
- [(c)](b) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
- 36. (currently amended) The isolated polypeptide of Claim 33 having at least 95% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[[)]; or
- (b) the amino acid sequence of the polypeptide shown in Figure 2 (SEQ ID NO:2), lacking its associated signal peptide; or
- [(c)](b) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
- 37. (currently amended) The isolated polypeptide of Claim 33 having at least 99% amino acid sequence identity to:
- (a) the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[[)]; or
- (b) the amino acid sequence of the polypeptide shown in Figure 2 (SEQ ID NO:2), lacking its associated signal peptide; or
- [(c)](b) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
 - 38. (currently amended) An isolated polypeptide comprising:

- (a) the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[[)]; or
- (b) the amino acid sequence of the polypeptide shown in Figure 2 (SEQ ID NO:2), lacking its associated signal peptide; or
- [(c)](b) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
- 39. (currently amended) The isolated polypeptide of Claim 38 comprising the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2[[)]].
- 40. (currently amended) The isolated polypeptide of Claim 38 comprising the amino acid sequence of the polypeptide shown in Figure 2 (of SEQ ID NO:2), lacking its associated signal peptide.
- 41. (previously presented) The isolated polypeptide of Claim 38 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203004.
- 42. (previously presented)A chimeric polypeptide comprising a polypeptide according to Claim 33 fused to a heterologous polypeptide.
- 43 (previously presented) The chimeric polypeptide of Claim 42, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.